## AMENDMENTS TO THE CLAIMS:

The listing of claims provided below will replace all prior versions and listings of claims in the above-captioned application:

## LISTING OF CLAIMS:

1-22. (Canceled)

- An isolated nucleic acid comprising a first nucleic acid having at least 80% homology to a reference nucleotide sequence wherein the reference sequence is wherein said first nucleic acid selectively hybridizes under stringent hybridization conditions to a second nucleic acid having a sequence selected from the group consisting of nucleotides 1-695 of SEQ ID NO:1, SEQ ID NO:1, and SEQ ID NO:2, and the complements thereof wherein said first nucleic acid encodes a protein essential for post-transcriptional inactivation, or encodes a portion of said protein.
- 24. (Canceled)
- 25. (Canceled)
- 26. (Canceled)
- 27. (Canceled)
- 28. (Currently amended)

  The isolated nucleic acid of claim 23 wherein said reference sequence is second nucleic acid has the sequence of nucleotides 1-695 of SEQ ID NO:1,

- 29. (Currently amended) The isolated nucleic acid of claim 23 wherein said reference sequence is second nucleic acid has a nucleotide sequence selected from the group consisting of SEQ ID NO:1 and SEQ ID NO:2.
- 30. (Canceled)
- 31. (Previously presented) An isolated nucleic acid comprising a nucleotide sequence having nucleotides 1-695 of SEQ ID NO:1.
- 32. (Previously presented) An isolated nucleic acid comprising a nucleotide sequence selected from the group consisting of SEQ ID NO:1 and SEQ ID NO:2.
- 33. (Previously presented) The isolated nucleic acid of claim 32 wherein said nucleotide sequence is SEQ ID NO:1.
- 34. (Previously presented) The isolated nucleic acid of claim 32 wherein said nucleotide sequence is SEQ ID NO:2.
- 35. (Previously presented) The isolated nucleic acid of claim 29 wherein said nucleic acid restores an sgs3 mutant of Arabidopsis thaliana.
- 36. (Withdrawn) An isolated polypeptide comprising an amino acid sequence having at least 80% homology to SEQ ID NO:3.
- 37. (Withdrawn) The isolated polypeptide of claim 36 wherein said amino acid sequence is at least 90% homologous to SEQ ID NO:3.

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- 38. (Withdrawn) The isolated polypeptide of claim 37 wherein said amino acid sequence is at least 95% homologous to SEQ ID NO:3.
- 39. (Withdrawn) The isolated polypeptide of claim 38 wherein said amino acid sequence is at least 98% homologous to SEQ ID NO:3.
- 40. (Withdrawn) The isolated polypeptide of claim 39 wherein said amino acid sequence is at least 99% homologous to SEQ ID NO:3.
- 41. (Withdrawn) The isolated polypeptide of claim 36 wherein said polypeptide restores an sgs3 mutant or Arabidopsis thaliana.
- 42. (Withdrawn) An isolated polypeptide comprising an amino acid sequence of SEQ ID NO:3.
- 43. (Withdrawn) An isolated polypeptide comprising a fragment of a polypeptide having an amino acid sequence of SEQ ID NO:3 wherein said fragment has biological activity in a plant or plant cell.
- 44. (Currently amended) An expression cassette comprising:
  a plant promoter;
  - a nucleic acid comprising a nucleotide sequence that is at least 80% homologous that
    selectively hybridizes under stringent hybridization conditions to SEQ ID NO:2
    and that encodes a protein essential for post-transcriptional inactivation, or
    encodes a portion of said protein; and

a plant terminator,

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wherein said plant promoter is operably linked to said nucleic acid, and wherein said terminator is operably linked to said nucleic acid.

- 45. (Currently amended) An expression cassette comprising:
  - a plant promoter;
  - a nucleic acid comprising a nucleotide sequence that is at least 80% homologous selectively hybridizes under stringent hybridization conditions to the complement of a nucleotide sequence selected from the group consisting of SEQ ID NO:1 and SEQ ID NO:2; and

a plant terminator,

wherein said plant promoter is operably linked to said nucleic acid, and wherein said terminator is operably linked to said nucleic acid.

- 46. (Canceled)
- 47. (Currently amended) An expression vector or transformation vector comprising a nucleic acid of claim 23, 28, or 29 or an expression cassette of claim 44, or 45, or 46.
- 48. (Currently amended) A process for transforming a <u>plant</u>, <u>yeast</u>, <u>flungal or bacterial</u> host organism comprising:
  - i. the contacting said plant, yeast, fungal or bacterial host organism with either a nucleic acid of claim 23, 28, or 29 or an expression cassette of claim 44, or 45, or

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- 46 under conditions that permit the internalization of said nucleic acid or said expression cassette into the host organism; and
- ii. selecting the host organism transformed with said nucleic acid or said expression cassette.
- 49. (Currently amended) A process for expressing a heterologous gene in a <u>plant</u>, <u>yeast, fungal or bacterial</u> host organism comprising:
  - heterologous gene, with an expression cassette comprising a plant promoter; a nucleic acid comprising a nucleotide sequence that is at least 80% homologous selectively hybridizes under stringent hybridization conditions to the complement of a nucleotide sequence selected from the group consisting of SEQ ID NO:1 and SEQ ID NO:2; and a plant terminator;
  - ii. selecting the host organism transformed with said expression cassette; and
     iii. culturing the selected host organism under conditions that permit the expression
    - of said nucleotide sequence contained within said expression cassette.
- 50. (Withdrawn) A process for expressing a heterologous gene in a host organism comprising contacting a host organism which comprises a heterologous gene, with a polypeptide comprising an amino acid sequence that is at least 80% homologous to SEQ ID NO:3.

- 51. (Currently amended) A transformed <u>plant, yeast, fungal or bacterial</u> host organism comprising at least one nucleic acid of claim 23, 28, or 29 or an expression cassette of claim 44, or 45, or 46.
- 52. (Currently amended) An isolated nucleic acid that selectively hybridizes <u>under stringent hybridization conditions</u> to a nucleic acid having a nucleotide sequence selected from the group consisting of the complement of nucleotides 1-695 of SEQ ID NO:1, the complement of SEQ ID NO:1, and the complement of SEQ ID NO:2, and the complements thereof.
- 53. (New) An isolated nucleic acid comprising a nucleotide sequence having at least 80% homology to a reference nucleotide sequence wherein said reference sequence is selected from the group consisting of SEQ ID NO:1 and SEQ ID NO:2.